

**EXHIBIT C3**

**BEST AVAILABLE COPY**

2d ET Am JRM EL 28

=> mhib Hinton

filled form  
12.30

2x 77 + 18 2.50

Cells trypsinized  
Buffer A remake => pH to 8

2N 1L 292 2.1 0.48 1.2  
2N 1L 353 2.8 0.86 0.9  
2N 1L 209 1.1 0.93 2.2

Buffer B remake => filled

2N 1L 235 1.4 0.7 1.75  
2N 1L 303 2.2 0.45 1.13

HPA: NP/DC (1.0/2.5)

Dilute  
5mm DTT, Prot, MSF,  
0.2mM vanadate

2N 1L 254 1.7 0.61 1.5  
2N 1L 281 2.0 0.51 1.37

2N 1L 270 2.0 0.51 1.28  
2N 1L 270 2.0 0.51 1.28

21 hwe 19  
- Forgot to vortex  
at 2N buffer A

2N 1L 300 2.2 0.46 1.15  
2N 1L 262 1.7 0.57 1.93

\* Sample 2N/8 got  
ice in 1.7

2N 1L 336 2.5 0.40 1.00  
2N 1L 255 1.7 0.60 1.50

2N 1L 273 2.0 0.50 1.26  
2N 1L 277 1.9 0.52 1.31

2N 1L 277 1.9 0.52 1.31  
2N 1L 217 1.2 0.82 2.01  
2N 1L 256 1.7 0.60 1.49

2N 1L 288 2.0 0.49 1.22  
2N 1L 257 1.7 0.54 1.48

2N 1L 303 2.2 0.45 1.13  
2N 1L 288 2.0 0.49 1.22

		(3) 1		(3) 5		(3) 10	
(1)	2N	2E	2N	2E	2N	2E	
	12	9	15	12.7	12.9	14.5	14.3
	(6)	(6)	(6)	(6)	(6)	(6)	
	1	1	5	10			
(2)	2N	2E	2N	2E	2N	2E	
	12.4	9	10.1	15	12.6	13.1	20.4
	(8)	(8)	(8)	(8)	(8)	(8)	
	1	1	5	10			
(3)	2N	2E	2N	2E	2N	2E	
	12	9	17.5	17.3	12.2	14.8	11.3
	(4)	(4)	(4)	(4)	(4)	(4)	
	1	1	5	10			
(4)	2N	2E	2N	2E	2N	2E	
	4.8	8.8	12	9	4.8+4.8	8.8+8.8	
	(M)	(M)	(M)	(M)	(M)	(M)	
	1	1	5	10			

Note: loaded 18 instead of 2.58 of 2E for ECAD gel 5 & prog  
loaded 0.48 instead of 18 2E for SCAT gel

Repeat

		(3) 1m		(3) 5m		(3) 10m	
(1)	2N	2E	2N	2E	2N	2E	
	4.8	8.8	6.1	5.1	5.1	5.1	2N=
	(6)	(6)	(6)	(6)	(6)	(6)	2E=
	1	1	5	10			
(2)	2N	2E	2N	2E	2N	2E	
	4.8	8.8	4.0	6.0	5.0	5.2	8.2
	(8)	(8)	(8)	(8)	(8)	(8)	
	1	1	5	10			
(3)	2N	2E	2N	2E	2N	2E	
	4.8	8.8	4.5	4.9	5.9	4.5	4.9
	(4)	(4)	(4)	(4)	(4)	(4)	
	1	1	5	10			
(4)	2N	2E	2N	2E	2N	2E	
	4.8	8.8	12	22			
	(M)	(M)	(M)	(M)	(M)	(M)	
	1	1	5	10			

Results:

Se Cell (4) / Set 1

β Cat: see small amt. of cleaved product not seen  
in utm. sample

p12<sup>own</sup>: definitely see Sh. ft, esp. in lower amt.  
regions

RCAT: no Sh. ft. but see ~ 2X decrease

Cells (1), (2), (3), Set 2

- No signal or very faint ⇒ Redox. c Pierce Regent

Cell (1) - DEND (3)

- control screened up, ~~22X~~ 22X
- 1nM (9), slight ↓ ECA
- 5nM (5), ↓ in ECA @ baseline, 3-4X ↓ in ECA by ET-1
- 10nM (3), ECA lower down to ~~undetectable~~ barely detectable

Cell (2) - VGTD (6)

- control screened up
- 1nM, 5nM: ECA levels similar between <sup>ET-1</sup> stimulated of unstimulated samples
- 10nM ~~screened~~ <sup>screened</sup> downing of <sup>ECA</sup> ET-1 in ET-1

Cell (3)

- control screened up
- 1nM: significant inhib. of ↓ ECA
- 1 & 9nM: Maximal downing of ECA by ET-1

Concl:

- too little protein + to short Ab inc. time
- control screened up 2° to quantitation error

⇒ requantitate sample

- load 25% extract
- 1" / 45" (1" / 2") no incubation

- YPAD, LGHD @ 10mM 5 effect on ECAD by ET-1

①

∴ Caspases 1, 4, 5, 9 not likely to be involved

⇒ do we have trial of ①, ② @ 1mM

- DEVD, IETD, and VEDD interfere to ↓ ECAD by ET-1.

- have already ruled at Caspases 3, 7

- have ruled in Caspase 8

- need to ✓ Caspases 6, 10.